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Table S17

S. sclerotiorum and B. cinerea orthologs of appressorium-associated genes in Magnaporthe oryzae.

S. sclerotiorum	B. cinerea B05.10	B. cinerea T4	M. oryzae	Function
SS1G_05586.1	BC1G_09439.1	BofuT4_P114010.1	pls1 (MGG_12594.6)	appressorium penetration; tetraspanin
No ortholog	No ortholog	No ortholog	mpg1 (MGG_10315.6)	adhesion; hydrophobin
No ortholog	No ortholog	No ortholog	pth11 (MGG_05871.6)	appressorium differentiation; integral membrane protein
SS1G_11636.1& SS1G_11637.1	BC1G_10952.1	BofuT4_P155910.1	pde1 (MGG_00111.6)	no penetration hyphae; aminophospholipid translocase (APT) family of P-type ATPase
SS1G_01851.1	BC1G_02517.1	BofuT4_P109590.1	emp1 (MGG_00527.6)	appressorium formation; extracellular matrix protein
SS1G_01602.1	BC1G_02467.1	BofuT4_P110100.1	atg8 (MGG_01062.6)	penetration hyphae; autophagy
SS1G_13339.1	BC1G_14516.1	BofuT4_P076990.1	pth2 (MGG_01721.6)	penetration hyphae; carnitine O-acetyl transferase
SS1G_07136.1	BC1G_10211.1	BofuT4_P086790.1	mst12 (MGG_12958.6)	appressorium maturation; steA transcription factor
SS1G_00637.1	BC1G_06518.1	BofuT4_P051820.1	chm1 (MGG_06320.6)	appressorium formation and penetration; PAK protein kinase
No ortholog	No ortholog	No ortholog	mas1 [#] (asg1;gas1) (MGG_12337.6)	appressorium penetration
SS1G_10311.1	BC1G_13581.1	No ortholog	mas2 [#] (MGG_04202.6)	appressorium penetration
	BC1G_08931.1 & BC1G_14711.1	BofuT4_P075230.1 & BofuT4_P017380.1	mas3 [#] (MGG_11595.6)	uncharacterized
SS1G_00173.1	BC1G_06114.1	BofuT4_P142900.1	mas3 [#] (MGG_00703.6)	uncharacterized
No ortholog	No ortholog	No ortholog	cas1 (MGG_09875.6)	uncharacterized
SS1G_14237.1	BC1G_12619.1	BofuT4_P142900.1	mas-related (MGG_00992.6)	uncharacterized
No ortholog	No ortholog	No ortholog	mas-related (MGG_02253.6)	uncharacterized

^{*}These genes are all annotated as "mas3" in the Broad genome site. The mas1 and mas2 designation were taken from Xue et al 2002. Plant Cell 14:2107-2119

Supplementary Table S17